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FIGURE 1

hRANTES	ASPYSSDT TPCCFAYIARPLPRAHIKEYFYTSKG	CSPNAVVFVTRNRQV	CANPEKKVREYINSLEHS
mMIP-1 $\alpha$	APYGADTPTACCFYS SRKIPRQFIVDYFETSSL	CSQPGVIFLTERNRQI	CADSKETWQEYITDLELNA
mMIP-1 $\beta$	APMGSDPFTSCCFYSYTSRQLHRSFVNDYYETSSL	CSKPAVVFLTKRGRQI	CANPSEPVTVEYMSDLELN
MCP-1	LAQPDAINAPVTCYNTNRKISVQRLASVRRITSSK	CPKEAVIFKTIIVAKEI	CADPKQKVQDSMDHLDKQTQTPKT
MCP-3	SPOGLAQPVGINSTTCCYRHFINKKIPKQRLSEYRRITSSH	CPREAVIFKTKLDKEI	CADFTQKVQDFMKHLDDKKTQTQPKL
CK $\beta$ -1	TKTESSSRGPHPECECFYTYTKIPRQRIMDYETNSQ	CSKPGIVFITXRGHSV	CTNPSDKVVDYIKDMK
CK $\beta$ -4	ASNFDCCLGYTDRILLHPKFIVGTRQLANBGCDINALIFHTKKLSV	CANPKQTWVKYIVRLLSKVKNN	CANPKQTWVKYIVRLLSKVKNN
CK $\beta$ -6	VVIPSPCCMFFVSKRIPENRNVVSQLSRST	CLKGGVIFTTKGQQF	CGDEKQEWQRYMKNLDAKQKKA
CK $\beta$ -7	AQVGYNKELCCLVYTSWQIFQKFIVDYSETSPQ	CFKPGVILLTKRGRQI	CADENKKRWVQK
CK $\beta$ -8	ENFVLLDRFHATSADCCISYTPRSIPCSLLSEYFETNSE	CSKPGVIFLTKGRRF	CANPSDKQVQVCMENLKLDTRIKTRKN
CK $\beta$ -9	SDAGGAQDCCLKYSQRKI PAKVVSRYRKQEPESLGCISI	PAILFLPRKRSQAEEL	CADPKELKVQQLMQHLCKTPSPQKPAQ
CK $\beta$ -10	ENPQGLAQPDALNVPSTCCPTFESSKKISLQRLKSYVITTSR	CPQKAVIFRTKLGEI	CADPEKAVQNYMKHLGRKAHTLKT
CK $\beta$ -11	PAPTLSGTNDAECCLSVTQKPIPGYIVNFHYLLKDGCRVP	PAVVFTTLRGRQL	CAPPDQPNVERIIQRLQRTSAKPKRRSS
CK $\beta$ -12	RSQPKVPEWNTPTSTCCCLKYTEKVLPRRLVVGTRKALN	CHLPALIIFVTRNREV	CTNPNDDWVQEYIKDNLPLLPTRNLST
CK $\beta$ -13	PYGANNMEDSVTCRDYVRYRLPLRVVVKHFYWTSDS	CPRPGVULLTFRDKBI	CADPRVPVWVKHILNKLSSQ
NAP-1/IL-8	SAKELRCQCIKITYSKPFPHPKFIKELRVIESGPHCANTEIIVKLSDGREL	CLDPKEVWVQRVVEKFLKRAENS	
NAP-2	AELRCMCIKITYSGIHPKNIQSLEVVIGKGTGHCNQVEVLATLKDGRKI	CLDDPADPRIKCIQVKKLAGDESAD	
hPF4	EAEEDGDLQCLCVKTTTSQVRPHITSLSEVIKAGPH	CPTAQLLATLKNRGI	CLDLQAPLYKKILKKLES
CK $\alpha$ -1	VLEVYTTSLRCRCVQESSVFIPRRFIDRIQILPRGNGCCPRKEIIVMKKNSIV	CVDPQAEWIQRMMEVLEKR	